

IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/001,407A

DATE: 08/27/2004 TIME: 11:18:38

Input Set : A:\GP117.SeqList.txt

Output Set: N:\CRF4\08272004\J001407A.raw

4 <110> APPLICANT: Yang, Yeasing
5 Burrell, Terrie
7 <120> TITLE OF INVENTION: Compositions and Methods for Detecting
8 Human Immunodeficiency Virus 2 (HIV-2)
11 <130> FILE REFERENCE: GP117-03.UT

C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/001,407A

C--> 13 <141> CURRENT FILING DATE: 2001-10-22

- 13 <150> PRIOR APPLICATION NUMBER: 60/242,620
- 14 <151> PRIOR FILING DATE: 2000-10-23
- 16 <150> PRIOR APPLICATION NUMBER: 60/280,058
- 17 <151> PRIOR FILING DATE: 2001-03-30
- 19 <160> NUMBER OF SEQ ID NOS: 34
- 21 <170> SOFTWARE: FastSEQ for Windows Version 3.0
- 23 <210> SEQ ID NO: 1
- 24 <211> LENGTH: 40
- 25 <212> TYPE: DNA
- 26 <213> ORGANISM: HIV-2
- 28 <400> SEQUENCE: 1
- 29 gtgtgtgttc ccatctctcc tagtcgccgc ctggtcattc 40
- 31 <210> SEQ ID NO: 2
- 32 <211> LENGTH: 19
- 33 <212> TYPE: DNA
- 34 <213> ORGANISM: HIV-2
- 36 <400> SEQUENCE: 2
- 37 gtgtgtgttc ccatctctc 19
- 39 <210> SEQ ID NO: 3
- 40 <211> LENGTH: 20
- 41 <212> TYPE: DNA
- 42 <213> ORGANISM: HIV-2
- 44 <400> SEQUENCE: 3
- 45 tgtgttccca tctctcctag 20
- 47 <210> SEQ ID NO: 4
- 48 <211> LENGTH: 21
- 49 <212> TYPE: DNA
- 50 <213> ORGANISM: HIV-2
- 52 <400> SEQUENCE: 4
- 53 gttcccatct ctcctagtcg c
- 55 <210> SEQ ID NO: 5
- 56 <211> LENGTH: 20
- 57 <212> TYPE: DNA
- 58 <213> ORGANISM: HIV-2
- 60 <400> SEQUENCE: 5
- 61 tcctagtcgc cgcctggtca



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63 <210> SEQ ID NO: 6 64 <211> LENGTH: 19 65 <212> TYPE: DNA 66 <213> ORGANISM: HIV-2 68 <400> SEQUENCE: 6 69 cctagtcgcc gcctqqtca 19 71 <210> SEQ ID NO: 7 72 <211> LENGTH: 20 73 <212> TYPE: DNA 74 <213> ORGANISM: HIV-2 76 <400> SEQUENCE: 7 77 tagtcgccgc ctggtcattc 20 79 <210> SEQ ID NO: 8 80 <211> LENGTH: 34 81 <212> TYPE: DNA 82 <213> ORGANISM: HIV-2 84 <400> SEQUENCE: 8 85 aaaatcccta gcaggttggc gcccgaacag ggac 34 87 <210> SEQ ID NO: 9 88 <211> LENGTH: 34 89 <212> TYPE: DNA 90 <213> ORGANISM: HIV-2 92 <400> SEQUENCE: 9 93 gtccctgttc gggcgccaac ctgctaggga tttt 34 95 <210> SEQ ID NO: 10 96 <211> LENGTH: 25 97 <212> TYPE: DNA 98 <213> ORGANISM: HIV-2 100 <400> SEQUENCE: 10 101 cgggcgccaa cctgctaggg atttt 25 103 <210> SEQ ID NO: 11 104 <211> LENGTH: 18 105 <212> TYPE: DNA 106 <213> ORGANISM: HIV-2 108 <400> SEQUENCE: 11 109 gtccctgttc gggcgcca 18 111 <210> SEQ ID NO: 12 112 <211> LENGTH: 23 113 <212> TYPE: DNA 114 <213> ORGANISM: Artificial Sequence 116 <220> FEATURE: 117 <223> OTHER INFORMATION: Primer mismatches HIV-2 sequence by deletion of 118 two nucleotides and mutation at a third position 120 <400> SEQUENCE: 12 121 cgggcgccac tgctagagat ttt 23 123 <210> SEQ ID NO: 13 124 <211> LENGTH: 24 125 <212> TYPE: DNA 126 <213> ORGANISM: Artificial Sequence

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128	<220>	FEATURE:	
129	<223>	OTHER INFORMATION: Primer mismatches HIV-2 sequence by deletion of	٠£
130		one nucleotide	Σ
132	<400>	SEQUENCE: 13	
133	cggg	cgccac ctgctaggga tttt	24
135	<210>	SEQ ID NO: 14	
136	<211>	LENGTH: 25	
137	<212>	TYPE: DNA	
		ORGANISM: HIV-2	
140	<400>	SEQUENCE: 14	
		gttcgg gcgccaacct gctag	25
		SEQ ID NO: 15	
		LENGTH: 52	
		TYPE: DNA	
146	<213>	ORGANISM: Artificial Sequence	
		FEATURE:	
149	<223>	OTHER INFORMATION: T7 promoter primer having a promoter sequence	
150		appended at the 5' end of an HIV-2 complementary	
151		primer sequence	
		SEQUENCE: 15	
154	aatt	taatac gactcactat agggagacgg gcgccaacct gctagggatt tt	52
		SEQ ID NO: 16	
		LENGTH: 45	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
162	<223>	OTHER INFORMATION: T7 promoter primer having a promoter sequence	
163 164		appended at the 5' end of an HIV-2 complementary	
		primer sequence SEQUENCE: 16	
169	<210>	caatac gactcactat agggagagte eetgtteggg egeca SEQ ID NO: 17	45
		LENGTH: 50	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: T7 promoter primer having a promoter sequence	
176		appended at the 5' end of the sequence given as	
177		SEQ ID NO:12	
179	<400>	SEQUENCE: 17	
		aatac gactcactat agggagacgg gcgccactgc tagagatttt	50
182	<210>	SEQ ID NO: 18	50
		LENGTH: 57	
184	<212>	TYPE: DNA	
185	<213>	ORGANISM: Artificial Sequence	
		FEATURE:	
188	<223>	OTHER INFORMATION: T7 promoter primer having a promoter sequence	
189		appended at the 5' end of the sequence given as	
190		SEQ ID NO:13	

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	<400> SEQUENCE: 18	
193	gaaattaata cgactcacta tagggagacc acacgggcgc cacctgctag ggatttt	57
195	<210> SEQ ID NO: 19	
196	<211> LENGTH: 58	
197	<212> TYPE: DNA	
198	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
201	<223> OTHER INFORMATION: T7 promoter primer having a promoter sequence	
202	appended at the 5' end of an HIV-2 complementary	
203		
205	<400> SEQUENCE: 19	
206	gaaattaata cgactcacta tagggagacc acaccctgtt cgggcgccaa cctgctag	58
208	<210> SEQ ID NO: 20	
209	<211> LENGTH: 22	
210	<212> TYPE: DNA	
211	<213> ORGANISM: HIV-2	
213	<400> SEQUENCE: 20	
	cetggtetgt taggaceett et	22
	<210> SEQ ID NO: 21	
217	<211> LENGTH: 14	
	<212> TYPE: DNA	
219	<213> ORGANISM: HIV-2	
221	<400> SEQUENCE: 21	
	gtctgttagg accc	14
224	<210> SEQ ID NO: 22	
	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213> ORGANISM: HIV-2	
	<400> SEQUENCE: 22	
	cctggtctgt taggaccc	18
	<210> SEQ ID NO: 23	
	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213> ORGANISM: HIV-2	
	<400> SEQUENCE: 23	
238	ctggtctgtt aggaccct	18
	<210> SEQ ID NO: 24	
_	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213> ORGANISM: HIV-2	
	<400> SEQUENCE: 24	
246	tggtctgtta ggaccctt	18
	<210> SEQ ID NO: 25	
	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213 > ORGANISM: HIV-2	
	<400> SEQUENCE: 25	
254	ggtctgttag gacccttc	18
∠ 56	<210> SEQ ID NO: 26	

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257	<211> LENGTH: 16						
258	<212> TYPE: DNA						
259	<213> ORGANISM: HIV-2						
261	<400> SEQUENCE: 26						
262	gtctgttagg accctt						16
264	<210> SEQ ID NO: 27			•			
265	<211> LENGTH: 18						
266	<212> TYPE: DNA						
267	<213> ORGANISM: HIV-2						
269	<400> SEQUENCE: 27						
270	gtctgttagg accettet						18
	<210> SEQ ID NO: 28						10
273	<211> LENGTH: 27						
274	<212> TYPE: DNA						
275	<213> ORGANISM: HIV-2						
277	<400> SEQUENCE: 28						
278	aatttaatac gactcactat	aqqqaqa					27
	<210> SEQ ID NO: 29	222 2					- /
281	<211> LENGTH: 33						
282	<212> TYPE: DNA						
283	<213> ORGANISM: HIV-2						
285	<400> SEQUENCE: 29						
286	gaaattaata cgactcacta	tagggagacc	aca				33
288	<210> SEQ ID NO: 30						
289	<211> LENGTH: 30						
290	<212> TYPE: RNA					-	
291	<213> ORGANISM: HIV-2						
293	<400> SEQUENCE: 30						
	gaaggguccu aacagaccag	ggucuuguua					30
296	<210> SEQ ID NO: 31					h _{rg}	
	<211> LENGTH: 53						
	<212> TYPE: DNA						
299	<213> ORGANISM: HIV-2						
301	<400> SEQUENCE: 31						
302	tteetgeege cettaetgee t	tttaaaaaaa	aaaaaaaaa	aaaaaaaaa	aaa		53
	<210> SEQ ID NO: 32						
	<211> LENGTH: 51						
	<212> TYPE: DNA	,					
307	<213> ORGANISM: HIV-2						
	<400> SEQUENCE: 32			•			
	ttcctgccgc ccttactgtt t	taaaaaaaaa	aaaaaaaaa	aaaaaaaaa	a		51
	<210> SEQ ID NO: 33						
	<211> LENGTH: 16						
	<212> TYPE: DNA						
	<213> ORGANISM: HIV-2						
	<400> SEQUENCE: 33						
	aaagggtcct aacaga						16
	<210> SEQ ID NO: 34						
321	<211> LENGTH: 9						

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/001,407A

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Input Set : A:\GP117.SeqList.txt

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date